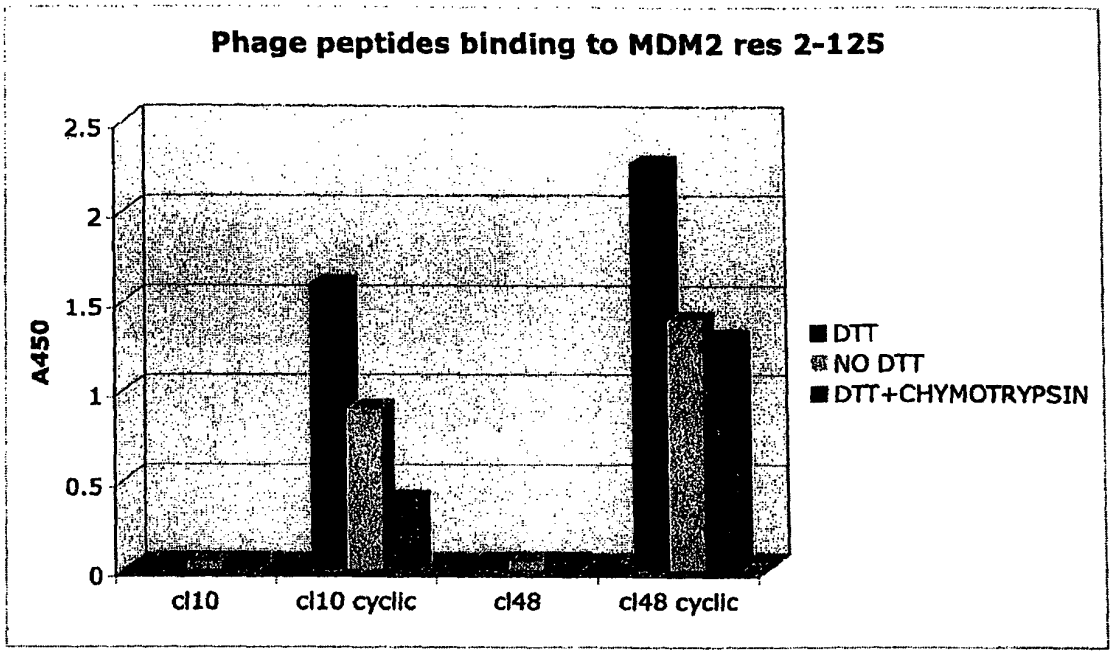

Phage peptides binding to MDM2 res 2-125
A450
2.5
2
1.5
1
0.5
0
cl10
cl10 cyclic
cl48
cl48 cyclic
DTT
NO DTT
DTT+CHYMOTRYPSIN

Figure 1

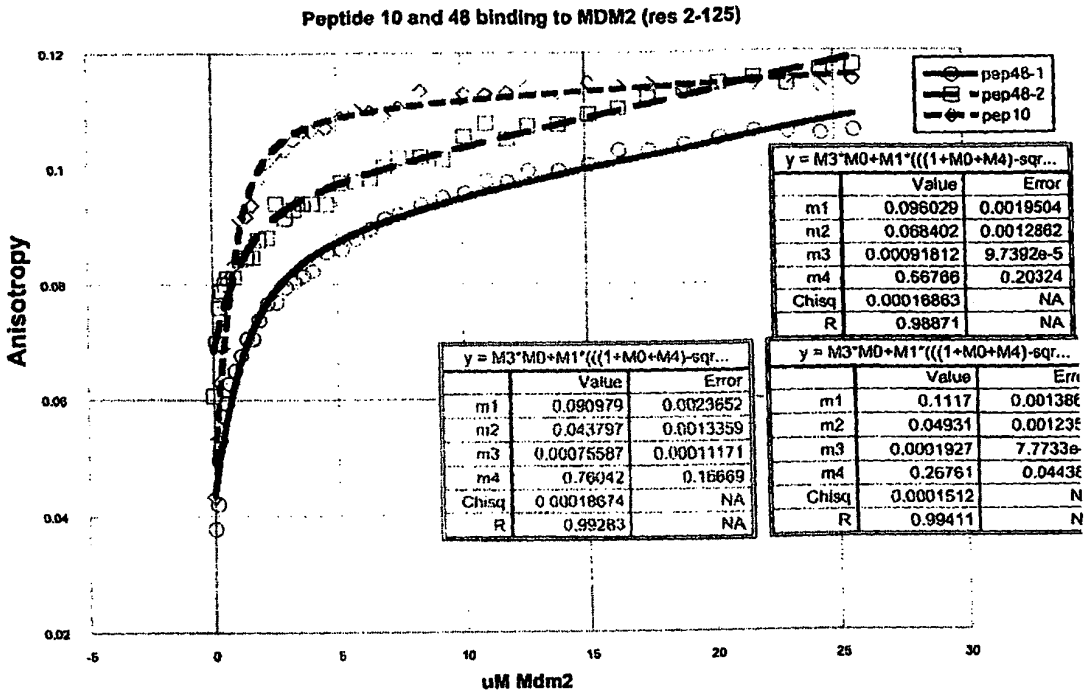

Peptide 10 and 48 binding to MDM2 (res 2-125)
Anisotropy
uM Mdm2
0.12
0.1
0.08
0.06
0.04
0.02
-5
0
5
10
15
20
25
30
pep48-1
pep48-2
pep10
y = M3*M0+M1*(((1+M0+M4)-sqr...
Value
Error
m1 0.096029 0.0019504
m2 0.068402 0.0012862
m3 0.00091812 9.7392e-5
m4 0.56786 0.20324
Chisq 0.00016863 NA
R 0.98871 NA
y = M3*M0+M1*(((1+M0+M4)-sqr...
Value
Error
m1 0.090979 0.0023652
m2 0.043797 0.0013359
m3 0.00075587 0.00011171
m4 0.76042 0.16669
Chisq 0.00018674 NA
R 0.99283 NA
y = M3*M0+M1*(((1+M0+M4)-sqr...
Value
m1 0.1117
m2 0.04931
m3 0.0001927
m4 0.26761
Chisq 0.0001512
R 0.99411

Figure 2